WO 2005/049093 PCT/EP2004/012421 1/26

SEQUENCE LISTING

														•	
<110>	INST	TTUT(១ នប	PERI	ORE	DI S	ANIT	A							
<120>	USE (OF M	ICRO	PART	ICLE	s FO	R AN'	TIGE	N DE	LIVE	RY				
<130>	N.89	060A	JHS		•										•
<160>	55														
<170>	Pate	ntIn	ver	sion	3.2										
<210> <211> <212> <213>	_	n im	muno	defi	ciend	cy v:	irus								
<220> <221> <222>	CDS (1).	. (30	9)												
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cag ccț Gln Pro															96
cat tgo His Cys															144
agg aag Arg Lys 50						_	_					_	_		192
cat caa His Gln 65															240
ccg aca Pro Thr															288
aca gat Thr Asp	_	_	_		tga										309
<211> <212>	2 102 PRT Human	ı imn	unoc	defic	cienc	y Vi	.rus								
<400>	2														
Met Glu 1	Pro	Val	Asp 5	Pro	Arg	Leu	Glu	Pro 10	Trp	Lys	His	Pro	Gly 15	Ser	

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 25 30

His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp 65 70 75 80

Pro Thr Gly Pro Lys Glu Gln Lys Lys Lys Val Glu Arg Glu Thr Glu 85 90 95

Thr Asp Pro Val His Gln 100

<210> 3

<211> 261

<212> DNA

<213> Human immunodeficiency virus

85

<220>

<221> CDS

<222> (1)..(261)

<400> 3

atg gag cca gta gat cct cgt cta gag ccc tgg aag cat cca gga agt

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser

1 10 15

cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt 96
Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
20 25 30

cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tac ggc
His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
35

agg aag aag cgg aga cag cgt cga aga cct cct caa ggc agt cag act 192
Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
50 55

cat caa gtt tct cta tca aag caa ccc acc tcc caa tcc cga ggg gac
His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
70 75 80

ccg aca ggc ccg aag gaa tag
261
Pro Thr Gly Pro Lys Glu

<210> <211> 86 <212> PRT <213> Human immunodeficiency virus <400> Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 1

10 15

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 30 25

His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 35 45 40

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr 55 60 50

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp 65 70 75 80

Pro Thr Gly Pro Lys Glu 85

<210> <211> 261 <212> DNA <213> Human immunodeficiency virus

<220> <221> CDS <222> (1)..(261)

<400> 5 atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 10 15

48

96 cag cct aaa act gct ggt acc aat tgc tat tgt aaa aag tgt tgc ttt Gln Pro Lys Thr Ala Gly Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 30

cat tgc caa gtt tgt ttc ata aca aaa gcc tta ggc atc tcc tat ggc 144 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 35 40 45

192 agg aag aag cgg aga cag cga cga aga cct cct caa ggc agt cag act Arg Lys Lys Arg Arg Gln Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 55 60

cat caa gtt tct cta tca aag cag ccc acc tcc caa tcc cga ggg gac 240 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp

WO 2005/049093		4.40.6	PCT/EP2004/012421
65	70	4/26 75	80
ccg aca ggc co Pro Thr Gly Pr			261
<210> 6 <211> 86 <212> PRT <213> Human 5	mmunodeficiency	virus	•
<400> 6			

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 1 5 15

Gln Pro Lys Thr Ala Gly Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 25 30

His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp 70 75 80

Pro Thr Gly Pro Lys Glu

<210> 7 <211> 261

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(261)

<400> 7

atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser

1 10 15

cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt 96
Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
20 25 30

cat tgc caa gtt tgt ttc ata aca gct gcc tta ggc atc tcc tat ggc
His Cys Gln Val Cys Phe Ile Thr Ala Ala Leu Gly Ile Ser Tyr Gly
35 40 45

WO 2005/049093										PCT/	EP20 0	4/0124	21					
							5/20	5										
agg	aad	aaq	caa	aga	caq	cqa	cga	aga	cct	cct	caa	ggc	agt	caq	act		192	

agg aag aag cgg aga cag cga cga aga cct cct caa ggc agt cag act
Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
50 55

cat caa gtt tct cta tca aag cag ccc acc tcc caa tcc cga ggg gac

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp

70

75

80

ccg aca ggc ccg aag gaa tag
Pro Thr Gly Pro Lys Glu
85

<210> 8 <211> 86 <212> PRT

<213> Human immunodeficiency virus

<400> 8

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 1 5 10 15

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 25 30

His Cys Gln Val Cys Phe Ile Thr Ala Ala Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp 65 70 75 80

Pro Thr Gly Pro Lys Glu 85

<210> 9 <211> 252 <212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(252)

<400> 9

atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser

1 10 15

cag cct aaa act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt 96
Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe

WO :	2005/	04909	3 20					6/20 25	5			30	PCT/	EP2004/012	421
		caa Gln 35													144
		aag Lys		_						_	_				192
		gtt Val				_	_					_			240
_	_	gaa Glu	tag												252
<210 <213 <212 <213	l> 2>	10 83 PRT Humar	n imr	nuno	defic	cienc	cy vi	irus					·		

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 1 5 10 15

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 20 25 30

His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Pro Thr Gly 65 70 75 80

Pro Lys Glu

<400>

10

<210> 11 <211> 252 <212> DNA <213> Human immunodeficiency virus

<220>
<221> CDS
<222> (1)..(252)

<400> 11

WO 2005/049093			PCT/EP2004/012421
		7/26	
		a gag ccc tgg aag 1 Glu Pro Trp Lys 10	
		tgc tat tgt aaa Cys Tyr Cys Lys 25	· · · · · · · · · · · · · · · · · · ·
		a gct gcc tta ggc Ala Ala Leu Gly	
		a aga cct cct caa g Arg Pro Pro Gln 60	
_	_	g ccc acc tcc caa n Pro Thr Ser Gln 75	
ccg aag gaa tag Pro Lys Glu			252
<210> 12 <211> 83 <212> PRT <213> Human imm <400> 12	unodeficiency v	virus	
Met Glu Pro Val	Asp Pro Arg Lev	Glu Pro Trp Lys	His Pro Gly Ser

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe

10

15

His Cys Gln Val Cys Phe Ile Thr Ala Ala Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Pro Pro Gln Gly Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Pro Thr Gly 70 75 80

Pro Lys Glu

<210> 13

<211> 306

<212> DNA

<213> Human immunodeficiency virus

<220> <221> CDS <222> (1)(30	06)	•		
			tgg aac cat ccg Trp Asn His Pro	
			tgt aaa aag tgt Cys Lys Lys Cys 30	
			tta ggc atc tcc Leu Gly Ile Ser 45	
			cct cag agc agt Pro Gln Ser Ser 60	
cat caa aat cct His Gln Asn Pro 65	ata cca aag Ile Pro Lys 70	caa ccc ata Gln Pro Ile	ccc caa acc caa Pro Gln Thr Gln 75	ggg gtc 240 Gly Val 80
tcg aca ggc ccg Ser Thr Gly Pro	gaa gaa tcg Glu Glu Ser 85	aag aag aag Lys Lys Lys 90	gtg gag agc aag Val Glu Ser Lys	gca gag 288 Ala Glu 95
aca gat cga ttc Thr Asp Arg Phe 100	Asp	•		306
<210> 14 <211> 101 <212> PRT <213> Human im	munodeficien	cy virus		•
<400> 14				•
Met Asp Pro Val 1	Asp Pro Asn 5	Leu Glu Pro 10	Trp Asn His Pro	Gly Ser 15
Gln Pro Thr Thr 20	Ala Cys Asn	Lys Cys Tyr 25	Cys Lys Lys Cys 30	Cys Tyr
His Cys Gln Val 35	Cys Phe Leu	Asn Lys Gly	Leu Gly Ile Ser 45	Tyr Gly
Arg Lys Lys Arg 50	Arg Gln Arg 55	Arg Gly Thr	Pro Gln Ser Ser 60	Lys Asp
His Gln Asn Pro 65	Ile Pro Lys 70	Gln Pro Ile	Pro_Gln Thr Gln 75	Gly Val 80

Ser Thr Gly Pro Glu Glu Ser Lys Lys Lys Val Glu Ser Lys Ala Glu 85 90 95

Thr Asp Arg Phe Asp 100

<210> 15

<211> 306

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(306)

<400> 15

atg gag cca gta gat cct aga cta gag ccc tgg aag cat cca gga agt

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser

1 10 15

cag cct aag act gct tgt acc aat tgc tat tgt aaa aag tgt tgc ttt 96
Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
20 25 30

cat tgc caa gtt tgt ttc ata aca aaa ggc tta ggc atc tcc tat ggc
His Cys Gln Val Cys Phe Ile Thr Lys Gly Leu Gly Ile Ser Tyr Gly
35
40
45

agg aag aag cgg aga cag cga cga aga gct cct caa gac agt cag act 192 Arg Lys Lys Arg Arg Gln Arg Arg Arg Ala Pro Gln Asp Ser Gln Thr 50 55

cat caa gtt tct cta tca aag caa ccc gcc tcc cag ccc cga ggg gac
His Gln Val Ser Leu Ser Lys Gln Pro Ala Ser Gln Pro Arg Gly Asp
70 75 80

ccg aca ggc ccg aag gaa tcg aag aag gtg gag aga gag aca gag
Pro Thr Gly Pro Lys Glu Ser Lys Lys Lys Val Glu Arg Glu Thr Glu
85
90
95

aca gat ccg gtc gat tag . 306
Thr Asp Pro Val Asp
100

<210> 16

<211> 101

<212> PRT

<213> Human immunodeficiency virus

<400> 16

Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser 1 5 10 15

Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe

PCT/EP2004/012421

10/26 20 25 30

His Cys Gln Val Cys Phe Ile Thr Lys Gly Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Gln Arg Arg Arg Ala Pro Gln Asp Ser Gln Thr 50 60

His Gln Val Ser Leu Ser Lys Gln Pro Ala Ser Gln Pro Arg Gly Asp 65 70 75 80

Pro Thr Gly Pro Lys Glu Ser Lys Lys Lys Val Glu Arg Glu Thr Glu 85 90 95

Thr Asp Pro Val Asp 100

<210> 17

<211> 306

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(306)

<400> 17

atg gag cca gta gat cct aac cta gag ccc tgg aac cat cca gga agt

Met Glu Pro Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly Ser

1 10 15

cag cct aaa act gct tgt aat aag tgt tat tgt aaa cac tgt agc tat 96
Gln Pro Lys Thr Ala Cys Asn Lys Cys Tyr Cys Lys His Cys Ser Tyr
20 25 30

cat tgt cta gtt tgc ttt cag aca aaa ggc tta ggc att tcc tat ggc
His Cys Leu Val Cys Phe Gln Thr Lys Gly Leu Gly Ile Ser Tyr Gly
35
40
45

agg aag aag cgg aga cag cga cga agc gct cct cca agc agt gag gat 192
Arg Lys Lys Arg Arg Gln Arg Arg Ser Ala Pro Pro Ser Ser Glu Asp
50 55

cat caa aat ctt ata tca aag caa ccc tta ccc caa acc caa ggg gac
His Gln Asn Leu Ile Ser Lys Gln Pro Leu Pro Gln Thr Gln Gly Asp
70 75 80

ccg aca ggc tcg gaa gaa tcg aag aag gtg gag agc aag aca gag
Pro Thr Gly Ser Glu Glu Ser Lys Lys Val Glu Ser Lys Thr Glu
85
90
95

aca gat cca ttc gat tag 306
Thr Asp Pro Phe Asp 100

<210> 18 <211> 101 <212> PRT Human immunodeficiency virus <213> <400> 18 Met Glu Pro Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly Ser 5 10 15 1 Gln Pro Lys Thr Ala Cys Asn Lys Cys Tyr Cys Lys His Cys Ser Tyr 20 25 30 His Cys Leu Val Cys Phe Gln Thr Lys Gly Leu Gly Ile Ser Tyr Gly 35 45 Arg Lys Lys Arg Arg Gln Arg Arg Ser Ala Pro Pro Ser Ser Glu Asp 50 55 60 His Gln Asn Leu Ile Ser Lys Gln Pro Leu Pro Gln Thr Gln Gly Asp 65 .70 75 80 Pro Thr Gly Ser Glu Glu Ser Lys Lys Lys Val Glu Ser Lys Thr Glu 85 95 90 Thr Asp Pro Phe Asp 100 <210> 19 <211> 261 <212> DNA <213> Human immunodeficiency virus <220> <221> CDS <222> (1)..(261) <400> 19 atg gat cca gta gat cct aac cta gag ccc tgg aac cat cca gga agt 48 Met Asp Pro Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly Ser 1 10 15 cag cct agg act cct tgt aac aag tgt tat tgt aaa aag tgt tgc tat 96 Gln Pro Arg Thr Pro Cys Asn Lys Cys Tyr Cys Lys Lys Cys Cys Tyr 20 25 30 cat tgc caa gtt tgc ttc ata acg aaa ggc tta ggc atc tcc tat ggc 144 His Cys Gln Val Cys Phe Ile Thr Lys Gly Leu Gly Ile Ser Tyr Gly 35 45 40 agg aag aag cgg aga cag cga cga aga cct cct caa ggc ggt cag gct 192

12/26

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Gly Gln Ala 50 55 60

cat caa gat cct ata cca aag caa ccc tcc tcc cag ccc cga ggg gac
His Gln Asp Pro Ile Pro Lys Gln Pro Ser Ser Gln Pro Arg Gly Asp
70 75 80

ccg aca ggc ccg aag gaa tag
Pro Thr Gly Pro Lys Glu
85

<210> 20 <211> 86 <212> PRT

<213> Human immunodeficiency virus

<400> 20

Met Asp Pro Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly Ser 1 5 10 15

Gln Pro Arg Thr Pro Cys Asn Lys Cys Tyr Cys Lys Lys Cys Cys Tyr 20 25 30

His Cys Gln Val Cys Phe Ile Thr Lys Gly Leu Gly Ile Ser Tyr Gly 35

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Gly Gln Ala 50 55 60

His Gln Asp Pro Ile Pro Lys Gln Pro Ser Ser Gln Pro Arg Gly Asp 65 70 75 80

Pro Thr Gly Pro Lys Glu 85

<210> 21 <211> 306

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(306)

<400> 21

atg gaa cta gta gat cct aac tta gat ccc tgg aac cat cca gga agc 48
Met Glu Leu Val Asp Pro Asn Leu Asp Pro Trp Asn His Pro Gly Ser
1 10 15

cag cct aca act cct tgt acc aaa tgc tat tgt aaa agg tgt tgc ttt 96
Gln Pro Thr Thr Pro Cys Thr Lys Cys Tyr Cys Lys Arg Cys Cys Phe
20 25 30

								13/2	6							
						aca Thr										144
agg Arg	aag Lys 50	aag Lys	cgg Arg	aga Arg	cag Gln	cga Arg 55	cga Arg	aga Arg	act Thr	cct Pro	caa Gln 60	agc Ser	agt Ser	cag Gln	ata Ile	192
		_		=		aag Lys								_		240
_			•			tcg Ser										288
	-	ccg Pro	_	gat Asp	tag											306
<210 <211 <212 <213	L> : 2> 1	22 LO1 PRT Humar	ı imr	nunoc	defic	cienc	cy vi	irus								
<400)> 2	22														
Met 1	Glu	Leu	Val	Asp 5	Pro	Asn	Leu	Asp	Pro 10	Trp	Asn	His	Pro	Gly 15	Ser	
Gln	Pro	Thr	Thr 20	Pro	Cys	Thr	Lys	Cys 25	Tyr	Cys	Lys	Arg	Cys 30	Cys	Phe	•
His	Cys	Gln 35	Trp	Cys	Phe	Thr	Thr 40	Lys	Gly	Leu	Gly	Ile 45	Ser	Tyr	Gly	
Arg	Lys 50	Lys	Arg	Arg	Gln	Arg 55	Arg	Arg	Thr	Pro	Gln 60	Ser	Ser	Gln	Ile	
His 65	Gln	Asp	Pro	Val	Pro 70	Lys	Gln	Pro	Leu	Ser 75	Gln	Ala	Arg	Gly	Asn 80	

Pro Thr Gly Pro Lys Glu Ser Lys Lys Glu Val Glu Ser Lys Ala Lys

90

95

Thr Asp Pro Cys Asp 100

<210> 23

<211> 306

<212> DNA

<213> Human immunodeficiency virus

85

<220> <221> CDS <222> (1)(306)	
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cag cct aaa act ccc tgt aac aaa tgt tat tgt aaa atg tgt tgc Gln Pro Lys Thr Pro Cys Asn Lys Cys Tyr Cys Lys Met Cys Cys 25 30	
cat tgt caa gtt tgc ttt ctg aac aaa ggc tta ggc atc tcc tat His Cys Gln Val Cys Phe Leu Asn Lys Gly Leu Gly Ile Ser Tyr 35 40 45	
agg aag aag cgg aag cac cga cga gga act cct cag agc agt aag Arg Lys Lys Arg Lys His Arg Arg Gly Thr Pro Gln Ser Ser Lys 50 55 60	
cat caa aat cct gta cca aag caa ccc tta ccc acc acc aga ggg His Gln Asn Pro Val Pro Lys Gln Pro Leu Pro Thr Thr Arg Gly 65 70 75	
ccg aca ggc ccg aag gaa tcg aag aag gag gtg gag agc aag aca Pro Thr Gly Pro Lys Glu Ser Lys Lys Glu Val Glu Ser Lys Thr 85 90 95	
aca gat cca ttc gat tag Thr Asp Pro Phe Asp 100	306
<pre> <210> 24 <211> 101 <212> PRT <213> Human immunodeficiency virus</pre>	-
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Met Asp Pro Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly 1 5	Ser
Gln Pro Lys Thr Pro Cys Asn Lys Cys Tyr Cys Lys Met Cys Cys 20 25 30	Trp
His Cys Gln Val Cys Phe Leu Asn Lys Gly Leu Gly Ile Ser Tyr 35 40 45	Gly
Arg Lys Lys Arg Lys His Arg Arg Gly Thr Pro Gln Ser Ser Lys 50 . 55 . 60	Asp
His Gln Asn Pro Val Pro Lys Gln Pro Leu Pro Thr Thr Arg Gly 65 70 75	Asn 80

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Pro Thr Gly Pro Lys Glu Ser Lys Lys Glu Val Glu Ser Lys Thr Glu 85 90 95

Thr Asp Pro Phe Asp 100

<210> 25

<211> 261

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(261)

<400> 25

atg gac cca gta gat cct aac caa gag ccc tgg aac cat cca gga agt

Met Asp Pro Val Asp Pro Asn Gln Glu Pro Trp Asn His Pro Gly Ser

1 10 15

cag cct aaa act gct tgt aac aat tgt tat tgt aaa aag tgc tgc tat 96
Gln Pro Lys Thr Ala Cys Asn Asn Cys Tyr Cys Lys Lys Cys Cys Tyr
20 25 30

cat tgc caa ttg tgc ttt tta aag aaa ggc tta ggc att tcc tat ggc
His Cys Gln Leu Cys Phe Leu Lys Lys Gly Leu Gly Ile Ser Tyr Gly
35
40
45

agg aag aag cgg agc cag cga cga gga act cct gca agt ttg caa gat

Arg Lys Lys Arg Ser Gln Arg Arg Gly Thr Pro Ala Ser Leu Gln Asp

50 55 60

cat caa aat cct ata cca aag caa ccc tta tcc cga acc cgc ggg gac
His Gln Asn Pro Ile Pro Lys Gln Pro Leu Ser Arg Thr Arg Gly Asp
70 75 80

ccg aca ggc ccg aag gaa tag
Pro Thr Gly Pro Lys Glu
85

<210> 26

<211> 86

<212> PRT

<213> Human immunodeficiency virus

<400> 26

Met Asp Pro Val Asp Pro Asn Gln Glu Pro Trp Asn His Pro Gly Ser 1 5 10 15

Gln Pro Lys Thr Ala Cys Asn Asn Cys Tyr Cys Lys Lys Cys Cys Tyr
20 25 30

16/26

His Cys	Gln	Leu	Cys	Phe	Leu	Lys	Lys	Gly	Leu	Gly	Ile	Ser	Tyr	Gly
	35					40					45			

Arg	Lys	Lys	Arg	Ser	Gln	Arg	Arg	Gly	Thr	Pro	Ala	Ser	Leu	Gln	Asp
	50					55					60				

His Gln Asn	Pro Ile	Pro Lys	Gln	Pro	Leu	Ser	Arg	Thr	Arg	Gly	Asp
65		70				75					80

Pro Thr Gly Pro Lys Glu 85

<210>	27		
<211>	306		
<212>	DNA		
<213>	Human	immunodeficiency	virus

<220>		
<221>	CDS	
<222>	(1).	. (306)

<400> 27

atg	gag	ctg	gta	gat	cct	aac	cta	gag	CCC	tgg	aat	cat	ccg	gga	agt	48
Met	Glu	Leu	Val	Asp	Pro	Asn	Leu	Glu	Pro	Trp	Asn	His	Pro	Gly	Ser	
1				5					10					15		

cag	cct	aca	act	gct	tgt	agc	aag	tgt	tac	tgt	aaa	ata	tgt	tgc	tgg	96	5
Gln	Pro	Thr	Thr	Ala	Cys	Ser	Lys	Cys	Tyr	Cys	Lys	Ile	Cys	Cys	Trp		
			20					25					30				

cat	tgc	caa	cta	tgc	ttt	ctg	aaa	aaa	ggc	tta	ggc	atc	tcc	tat	ggc	144
His	Cys	Gln	Leu	Cys	Phe	Leu	Lys	Lys	Gly	Leu	Gly	Ile	Ser	Tyr	Gly	
		35					40					45				

agg	aag	aag	cgg	aaq	cac	cga	cga	gga	act	cct	caq	agc	aqt	aaq	gat	192
	=			_	His		_	_					_	_	=	
AT G	пуз	пуз	ALG	T) S	HTS	AT A	ALG	GTA	T 11T	FIO	GIII	Ser	Ser	പുട	vob	
	50					55					60					

cat	caa	aat	cct	ata	cca	നമന	caa	CCC	cta	CCC	atc	atc	ana	aaa	aac	240
													_		Asn	240
65	U				70					75	220		9	- 1	80	

			gaa Glu							288
		85			90			95	•	

|--|

<210> 28 <211> 101 <212> PRT <213> Human immunodeficiency virus <400> 28

Met Glu Leu Val Asp Pro Asn Leu Glu Pro Trp Asn His Pro Gly Ser 1 5 10 15

Gln Pro Thr Thr Ala Cys Ser Lys Cys Tyr Cys Lys Ile Cys Cys Trp
20 25 30

His Cys Gln Leu Cys Phe Leu Lys Lys Gly Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Lys His Arg Arg Gly Thr Pro Gln Ser Ser Lys Asp 50 60

His Gln Asn Pro Ile Pro Glu Gln Pro Leu Pro Ile Ile Arg Gly Asn 65 70 75 80

Pro Thr Asp Pro Lys Glu Ser Lys Lys Glu Val Ala Ser Lys Ala Glu 85 90 95

Thr Asp Pro Cys Asp 100

<210> 29

<211> 306

<212> DNA

<213> Human immunodeficiency virus

<220>

<221> CDS

<222> (1)..(306)

<400> 29

atg gag ccg gta gat cct agc cta gag ccc tgg aac cac ccg gga agt

Met Glu Pro Val Asp Pro Ser Leu Glu Pro Trp Asn His Pro Gly Ser

1 10 15

cag cct aca act gct tgt agc aat tgt tac tgt aaa atg tgc tgc tgg 96
Gln Pro Thr Thr Ala Cys Ser Asn Cys Tyr Cys Lys Met Cys Cys Trp
20 25 30

cat tgc caa ttg tgc ttt ctg aac aag ggc tta ggc atc tcc tat ggc
His Cys Gln Leu Cys Phe Leu Asn Lys Gly Leu Gly Ile Ser Tyr Gly
35
40
45

agg aag aag cgg aga cgc cga cga gga act cct cag agc cgt cag gat 192
Arg Lys Lys Arg Arg Arg Arg Gly Thr Pro Gln Ser Arg Gln Asp
50 55

cat caa aat cct gta cca aag caa ccc tta ccc acc acc aga ggg aac
His Gln Asn Pro Val Pro Lys Gln Pro Leu Pro Thr Thr Arg Gly Asn
70 75 80

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								18/2	6							
ccg	aca	ggc	ccg	aaa	gaa	tcg	aag	aag	gag	gtg	gcg	agc	aag	aca	gag	288

ccg aca ggc ccg aaa gaa tcg aag aag gag gtg gcg agc aag aca gag Pro Thr Gly Pro Lys Glu Ser Lys Lys Glu Val Ala Ser Lys Thr Glu 85 90 95

aca gat ccg tgc gat tag
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His Cys Gln Leu Cys Phe Leu Asn Lys Gly Leu Gly Ile Ser Tyr Gly 35 40 45

Arg Lys Lys Arg Arg Arg Arg Gly Thr Pro Gln Ser Arg Gln Asp 50 55 60

His Gln Asn Pro Val Pro Lys Gln Pro Leu Pro Thr Thr Arg Gly Asn 65 70 75 80

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WO 2005	04909	3					19/2	26					PCT	EP2004/012	2421
		20					25					30			
cat tgc His Cys															144
agg aag Arg Lys 50															192
caa gat Gln Asp 65		_													24.0
aaa cgc Lys Arg		_	_	_			_				_				288
ggt gga Gly Gly			_		_	_		_	_	_	_				336
tca gga Ser Gly		taa													348
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His Cys	Tyr 35	Val	Cys	Phe	Ala	Ser 40	Lys	Gly	Leu	Gly	Ile 45	Ser	Tyr	Gly	
Arg Lys 50	Lys	Arg	Arg	Arg	Pro 55	Ala	Ala	Ala	Ala	Ser 60	His	Pro	Asp	Asn	

Gln Asp Pro Val Pro Glu Gln Pro Pro Ser Ile Thr Asn Arg Lys Gln 65 70 75 80

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Gly Gly Tyr Pro Arg Arg Lys Asp Ser Cys His_Cys Cys Thr Arg Thr 100 105 110

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22/26

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Ser Thr Arg Thr Gln Ile Asn Lys Val Val Arg Phe Asp Lys Leu Pro 50 55 60

Gly Phe Gly Asp Ser Ile Glu Ala Gln Cys Gly Thr Ser Val Asn Val 65 70 75 80

His Ser Ser Leu Arg Asp Ile Leu Asn Gln Ile Thr Lys Pro Asn Asp 90 95

Val Tyr Ser Phe Ser Leu Ala Ser Arg Leu Tyr Ala Glu Glu Arg Tyr 100 105 110

Pro Ile Leu Pro Glu Tyr Leu Gln Cys Val Lys Glu Leu Tyr Arg Gly 115 120 125

Gly Leu Glu Pro Ile Asn Phe Gln Thr Ala Ala Asp Gln Ala Arg Glu 130 135 140

Leu Ile Asn Ser Trp Val Glu Ser Gln Thr Asn Gly Ile Ile Arg Asn 145 150 155 160

Val Leu Gln Pro Ser Ser Val Asp Ser Gln Thr Ala Met Val Leu Val 165 170 175

Asn Ala Ile Val Phe Lys Gly Leu Trp Glu Lys Ala Phe Lys Asp Glu 180 185 190

Asp Thr Gln Ala Met Pro Phe Arg Val Thr Glu Gln Glu Ser Lys Pro 195 200 205

Val Gln Met Met Tyr Gln Ile Gly Leu Phe Arg Val Ala Ser Met Ala 210 -220

24.	/26
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Ser Met Leu Val Leu Leu Pro Asp Glu Val Ser Gly Leu Glu Gln Leu 245 250 255

Glu Ser Ile Ile Asn Phe Glu Lys Leu Thr Glu Trp Thr Ser Ser Asn 260 265 270

Val Met Glu Glu Arg Lys Ile Lys Val Tyr Leu Pro Arg Met Lys Met 275 280 285

Glu Glu Lys Tyr Asn Leu Thr Ser Val Leu Met Ala Met Gly Ile Thr 290 295 300

Asp Val Phe Ser Ser Ser Ala Asn Leu Ser Gly Ile Ser Ser Ala Glu 305 310 315 320

Ser Leu Lys Ile Ser Gln Ala Val His Ala Ala His Ala Glu Ile Asn 325 330 335

Glu Ala Gly Arg Glu Val Val Gly Ser Ala Glu Ala Gly Val Asp Ala 340 345 350

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